

RAW SEQUENCE LISTING

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Application Serial Number: 10/594, 014
Source: TFWP
Date Processed by STIC: 10/3/86

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IFWP

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/594,014

DATE: 10/03/2006

TIME: 09:09:35

Input Set : A:\PTO.RJ.txt

Output Set: N:\CRF4\10032006\J594014.raw

3 <110> APPLICANT: DSM IP Assets B.V.
 4 Petrus J.T. Dekker
 5 Marco A. van den Berg
 7 <120> TITLE OF INVENTION: FILAMENTOUS FUNGAL MUTANTS WITH IMPROVED HOMOLOGOUS
 RECOMBINATION
 8 EFFICIENCY
 10 <130> FILE REFERENCE: 3663-335 / 24181USWO
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/594,014
 13 <141> CURRENT FILING DATE: 2006-09-25
 15 <150> PRIOR APPLICATION NUMBER: PCT/EP2005/051464
 16 <151> PRIOR FILING DATE: 2005-03-31
 18 <150> PRIOR APPLICATION NUMBER: EP 04076057.1
 19 <151> PRIOR FILING DATE: 2004-04-02
 21 <160> NUMBER OF SEQ ID NOS: 23
 23 <170> SOFTWARE: MS Word
 25 <210> SEQ ID NO: 1
 26 <211> LENGTH: 2284
 27 <212> TYPE: DNA
 28 <213> ORGANISM: Aspergillus niger
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 33 cagagctaca aaccagtcaa agatgcggtc ctcttcgcaa tcgatgtcag cgattccatg 180
 34 ttgacgcctc gcccctcagc agatcctaag aaacacaccc aagaatcacc caccacggca 240
 35 ggcgtcaaat ggcctatca cttcatgcaa caacgaatca tatcaaattc acaagacatg 300
 36 atgggtgttt tgctgtcgg gacccaggcg tccaaatttct ttgaagaaga tgaagacagt 360
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 38 catgaggtca aaggacttcg agcaactggta gatgatgaag gagactcaag ggaggttcta 480
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 49 gagtgacttc ccactccgca gatgatactg gcccggactgt cgagaaatgg gagatcagaa 1140
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57	tgattgatca	gatgcgcgtg	atcgccagc	aactgcagct	gccgaaggga	gtgtacgagc	1620
58	ctctcaaata	ccccatcca	tgtaaagtac	ttctgtcttg	cattgctcg	atacgatgaa	1680
59	cgagaagctg	acagccgtg	atcagccctt	caatgcatt	accgcattct	acaagctctc	1740
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62	ctgacaattg	caccacagcg	cgccgggtgac	tacgtattat	cctggggccg	cgaactcgaa	1920
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98	tttccgcatt	tgcatcagac	tcttttgcgt	tccaaagaaga	tggactctgt	ctgggttcatt	1260
99	gcacgcaagg	gtgctggccc	cggtctcgcc	gctatgatcg	caggcgaaga	aaagcttgc	1320
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104	ctccccgaaa	aaccagaaga	aaaaccatt	ccgaaatacc	gccaaatcga	caagcgcgc	1620
105	ggtgactacg	tattatcctg	ggccgacgaa	ctcgaaaagc	aatacgc	aacctcagca	1680
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 126 35 40 45
 128 Asp Pro Lys Lys His Thr Gln Glu Ser Pro Thr Thr Ala Ala Leu Lys
 129 50 55 60
 131 Cys Ala Tyr His Phe Met Gln Gln Arg Ile Ile Ser Asn Pro Gln Asp
 132 65 70 75 80
 134 Met Met Gly Val Leu Leu Phe Gly Thr Gln Ala Ser Lys Phe Phe Glu
 135 85 90 95
 137 Glu Asp Glu Asp Ser Arg Gly Asp Leu Ser Tyr Pro Asn Cys Tyr Leu
 138 100 105 110
 140 Phe Thr Asp Leu Asp Val Pro Ser Ala His Glu Val Lys Gly Leu Arg
 141 115 120 125
 143 Ala Leu Val Asp Asp Glu Gly Asp Ser Arg Glu Val Leu Ser Pro Ala
 144 130 135 140
 146 Lys Glu Gln Val Ser Met Ala Asn Val Leu Phe Cys Ala Asn Gln Ile
 147 145 150 155 160
 149 Phe Thr Ser Arg Ala Pro Asn Phe Leu Ser Arg Arg Leu Phe Ile Ile
 150 165 170 175
 152 Thr Asp Asn Asp Asn Pro His Gly Asp Asp Lys Thr Leu Arg Ser Ala
 153 180 185 190
 155 Ala Thr Val Arg Ala Lys Asp Leu Tyr Asp Leu Gly Val Thr Ile Glu
 156 195 200 205
 158 Leu Phe Pro Ile Ser Arg Pro Glu His Glu Phe Lys Asn Ser Lys Phe
 159 210 215 220
 161 Tyr Asp Ser Leu Pro Ser Asp Pro Glu Ala Pro Ala Tyr Leu Gln Ser
 162 225 230 235 240
 164 Asp Ser Lys Ala Ala Thr Ala Thr Gly Asp Gly Ile Ser Leu Leu Asn
 165 245 250 255
 167 Thr Leu Leu Ser Ser Ile Asn Ser Arg Thr Val Pro Arg Arg Thr His
 168 260 265 270
 170 Phe Ser Asn Met Pro Leu Glu Leu Gly Pro Asp Phe Arg Ile Ser Val
 171 275 280 285
 173 Ser Gly Tyr Ile Leu Leu Arg Arg Gln Ala Pro Ala Arg Asn Ser Phe
 174 290 295 300
 176 Ile Trp Leu Asn Gly Glu Lys Pro Val Val Ala Lys Gly Val Thr Ser
 177 305 310 315 320

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 180 325 330 335
 182 Lys Ala Tyr Lys Phe Gly Gly Asp Gln Val Thr Phe Ser Pro Asp Glu
 183 340 345 350
 185 Gln Lys Ala Leu Arg Asp Phe Gly Glu Pro Val Ile Arg Val Ile Gly
 186 355 360 365
 188 Phe Lys Pro Ile Thr Ala Leu Pro Phe Trp Ala Asn Val Lys His Pro
 189 370 375 380
 191 Tyr Phe Ile Tyr Pro Ser Glu Glu Asp Tyr Val Gly Ser Ser Arg Val
 192 385 390 395 400
 194 Phe Ser Ala Leu His Gln Thr Leu Leu Arg Ser Lys Lys Met Ala Leu
 195 405 410 415
 197 Val Trp Phe Ile Ala Arg Lys Gly Ala Gly Pro Val Leu Ala Ala Met
 198 420 425 430
 200 Ile Ala Gly Glu Glu Lys Leu Asp Glu Asn Gly Val Gln Lys Tyr Pro
 201 435 440 445
 203 Pro Gly Met Trp Ile Leu Pro Leu Pro Phe Ala Asp Asp Ile Arg Gln
 204 450 455 460
 206 Asn Pro Glu Thr Thr Leu Asn Val Ala Pro Glu Ser Leu Ile Asp Gln
 207 465 470 475 480
 209 Met Arg Val Ile Val Gln Gln Leu Gln Leu Pro Lys Gly Val Tyr Glu
 210 485 490 495
 212 Pro Leu Lys Tyr Pro Asn Pro Ser Leu Gln Trp His Tyr Arg Ile Leu
 213 500 505 510
 215 Gln Ala Leu Ala Leu Asp Glu Asp Leu Pro Glu Lys Pro Glu Asp Lys
 216 515 520 525
 218 Thr Ile Pro Lys Tyr Arg Gln Ile Asp Lys Arg Ala Gly Asp Tyr Val
 219 530 535 540
 221 Leu Ser Trp Ala Asp Glu Leu Glu Lys Gln Tyr Ala Lys Thr Ser Ala
 222 545 550 555 560
 224 Ala Ala Pro Arg Pro Thr Ser Thr Leu Val Lys Arg Gly Ser Lys Asp
 225 565 570 575
 227 Arg Ala Ser Glu Thr Glu Asp Ser Lys Pro Ser Lys Lys Ile Lys Val
 228 580 585 590
 230 Glu Glu Asp Ser Gly Ser Leu Glu Glu Glu Val Arg Arg His His Lys
 231 595 600 605
 233 Lys Gly Thr Leu Ser Lys Leu Thr Val Ala Ile Leu Lys Asp Phe Leu
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L:12 M:270 C: Current Application Number differs, Wrong Format